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RAW SEQUENCE LISTING

DATE: 03/19/2002 TIME: 16:30:50

PATENT APPLICATION: US/10/090,466

Input Set : A:\LEX-0315-USA SEQLIST.txt Output Set: N:\CRF3\03192002\J090466.raw

- 4 <110> APPLICANT: Turner, C. Alexander Jr.
- Mathur, Brian 5
- Cullinan, Emily B.
- 8 <120> TITLE OF INVENTION: Novel Human Dectin Proteins and Polynucleotides Encoding the

Same

- 10 <130> FILE REFERENCE: LEX-0315-USA
- C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/090,466
- C--> 12 <141> CURRENT FILING DATE: 2002-03-01
 - 12 <150> PRIOR APPLICATION NUMBER: US 60/274,961
 - 13 <151> PRIOR FILING DATE: 2001-03-12
 - 15 <160> NUMBER OF SEQ ID NOS: 4
 - 17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 - 19 <210> SEQ ID NO: 1
 - 20 <211> LENGTH: 642
 - 21 <212> TYPE: DNA
 - 22 <213> ORGANISM: homo sapiens
 - 24 <400> SEQUENCE: 1
 - 25 atqqtqcctq aaqaaqaqcc tcaaqaccqa qaqaaaqqac tctgqtqqtt ccaqttqaag 60
 - 26 gtctggtcca tggcagtcgt atccatcttg ctcctcagtg tctgtttcac tgtgagttct 120
 - 27 gtqgtgcctc acaattttat gtatagcaaa actgtcaaga ggctgtccaa gttacgagag 180
 - 28 tatcaacagt atcatccaag cctgacctgc gtcatggaag gaaaggacat agaagattgg 240
 - 29 agetgetgee caacceettg gaetteattt cagtetagtt getaetttat ttetaetggg 300
 - 30 atgeaatett ggaetaagag teaaaagaae tgttetgtga tgggggetga tetggtggtg 360
 - 31 atcaacacca gggaagaaca ggatttcatc attcagaatc tgaaaagaaa ttcttcttat 420
 - 32 tttctggggc tgtcagatcc agggggtcgg cgacattggc aatgggttga ccagacacca 480
 - 33 tacaatgaaa atgtcacatt ctggcactca ggtgaaccca ataaccttga tgagcgttgt 540

 - 34 qcqataataa atttccqttc ttcaqaaqaa tggggctgga atgacattca ctgtcatgta 600 35 cctcagaagt caatttgcaa gatgaagaag atctacatat aa
 - 37 <210> SEQ ID NO: 2
 - 38 <211> LENGTH: 213
 - 39 <212> TYPE: PRT
 - 40 <213> ORGANISM: homo sapiens
 - 42 <400> SEQUENCE: 2
 - 43 Met Val Pro Glu Glu Glu Pro Gln Asp Arg Glu Lys Gly Leu Trp Trp
 - 5 44 1 10
 - 45 Phe Gln Leu Lys Val Trp Ser Met Ala Val Val Ser Ile Leu Leu Leu
 - 46 20
 - 47 Ser Val Cys Phe Thr Val Ser Ser Val Val Pro His Asn Phe Met Tyr
 - 48
 - 49 Ser Lys Thr Val Lys Arg Leu Ser Lys Leu Arg Glu Tyr Gln Gln Tyr
 - 50
 - 51 His Pro Ser Leu Thr Cys Val Met Glu Gly Lys Asp Ile Glu Asp Trp
 - 52 65 70
 - 53 Ser Cys Cys Pro Thr Pro Trp Thr Ser Phe Gln Ser Ser Cys Tyr Phe



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90 55 Ile Ser Thr Gly Met Gln Ser Trp Thr Lys Ser Gln Lys Asn Cys Ser 105 57 Val Met Gly Ala Asp Leu Val Val Ile Asn Thr Arg Glu Glu Gln Asp 58 115 120 59 Phe Ile Ile Gln Asn Leu Lys Arg Asn Ser Ser Tyr Phe Leu Gly Leu 135 61 Ser Asp Pro Gly Gly Arg Arg His Trp Gln Trp Val Asp Gln Thr Pro 150 155 63 Tyr Asn Glu Asn Val Thr Phe Trp His Ser Gly Glu Pro Asn Asn Leu 64 65 Asp Glu Arq Cys Ala Ile Ile Asn Phe Arq Ser Ser Glu Glu Trp Gly 66 180 185 67 Trp Asn Asp Ile His Cys His Val Pro Gln Lys Ser Ile Cys Lys Met 68 195 200 69 Lys Lys Ile Tyr Ile 70 210 73 <210> SEQ ID NO: 3 74 <211> LENGTH: 549 75 <212> TYPE: DNA 76 <213> ORGANISM: homo sapiens 78 <400> SEQUENCE: 3 79 atggtgcctg aagaagagcc tcaagaccga gtgcctcaca attttatgta tagcaaaact 60 80 gtcaagaggc tgtccaagtt acgagagtat caacagtatc atycaagcct gacctgcgtc 120 81 atggaaggaa aggacataga agattggagc tgctgcccaa ccccttggac ttcatttcag 180 82 totagttgct actttatttc tactgggatg caatcttgga ctaagagtca aaagaactgt 240 83 totgtgatgg gggctgatot ggtggtgato aacaccaggg aagaacagga titcatcatt 300 84 cagaatctga aaagaaattc ttcttatttt ctggggctgt cagatccagg gggtcggcga 360 85 cattggcaat gggttgacca gacaccatac aatgaaaatg tcacattctg gcactcaggt 420 86 gaacccaata accttgatga gcgttgtgcg ataataaatt tccgttcttc agaagaatgq 480 87 ggctggaatg acattcactg tcatgtacct cagaagtcaa tttgcaagat gaagaagatc 540 88 tacatataa 90 <210> SEQ ID NO: 4 91 <211> LENGTH: 182 92 <212> TYPE: PRT 93 <213> ORGANISM: homo sapiens 95 <220> FEATURE: 96 <221> NAME/KEY: VARIANT 97 <222> LOCATION: (1)...(182) 98 <223> OTHER INFORMATION: Xaa = Any Amino Acid 100 <400> SEQUENCE: 4 101 Met Val Pro Glu Glu Glu Pro Gln Asp Arg Val Pro His Asn Phe Met 102 1 103 Tyr Ser Lys Thr Val Lys Arg Leu Ser Lys Leu Arg Glu Tyr Gln Gln W--> 105 Tyr His Xaa Ser Leu Thr Cys Val Met Glu Gly Lys Asp Ile Glu Asp 107 Trp Ser Cys Cys Pro Thr Pro Trp Thr Ser Phe Gln Ser Ser Cys Tyr

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109 110	Phe 65	Ile	Ser	Thr	Gly	Met 70	Gln	ser	Trp	Thr	Lys 75	Ser	Gln	Lys	Asn	Cys 80
111 112	Ser	Val	Met	Gly	Ala 85	Asp	Leu	Val	Val	Ile 90	Asn	Thr	Arg	Glu	Glu 95	Gln
113 114	Asp	Phe	Ile	Ile 100	Gln	Asn	Leu	Lys	Arg 105	Asn	Ser	Ser	Tyr	Phe 110	Leu	Gly
115 116	Leu	Ser	Asp 115	Pro	Gly	Gly	Arg	Arg 120	His	Trp	G1n	Trp	Val 125	Asp	Gln	Thr
117 118	Pro	Tyr 130	Asn	Glu	Asn	Val	Thr 135	Phe	Trp	His	ser	Gly 140	Glu	Pro	Asn	Asn
	Leu 145	Asp	Glu	Arg	Cys	Ala 150	Ile	Ile	Asn	Phe	Arg 155	Ser	Ser	Glu	Glu	Trp 160
121 122	Gly	Trp	Asn	Asp	11e 165	His	Cys	His	Va1	Pro 170	Gln	Lys	Ser	Ile	Cys 175	Lys
123 124	Met	Lys	Lys	Ile 180	Tyr	Ile										

VERIFICATION SUMMARY

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Input Set : A:\LEX-0315-USA SEQLIST.txt
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L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:105 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4